In the Claims:

1-118. Canceled.

- 119. (Currently amended) An isolated nucleic acid having at least 80% nucleic acid sequence identity to:
- (a) a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418);
- (b) a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418), lacking its associated signal peptide;
- (c) a nucleic acid sequence encoding the extracellular domain of the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418);
- (d) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in Figure 300 (SEQ ID NO:418), lacking its associated signal peptide;
- (e)(d) the nucleic acid sequence of SEQ ID NO: 417 shown in Figure 299 (SEQ ID NO:417);
- (f)(e) the full-length coding sequence of the nucleic acid sequence of SEQ ID NO: 417 shown in Figure 299 (SEQ ID NO:417); or
- (g)(f) the full-length coding sequence of the cDNA deposited under ATCC accession number 203115;

 wherein said polypeptide induces proliferation of stimulated T lymphocytes in a mixed lymphocyte reaction.
- 120. (Currently amended) An isolated nucleic acid of Claim 119 having at least 85% nucleic acid sequence identity to:
- (a) a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418);
- (b) a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418), lacking its associated signal peptide;
- (c) a nucleic acid sequence encoding the extracellular domain of the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418);

- (d) a nucleic acid-sequence encoding the extracellular domain of the polypeptide shown in Figure 300 (SEQ ID NO:418), lacking its associated signal peptide;
- (e) the nucleic acid sequence of SEQ ID NO: 417 shown in Figure 299 (SEQ ID NO:417);
- (f)(d) the full-length coding sequence of the nucleic acid sequence of SEQ ID NO: 417 shown in Figure 299 (SEQ ID NO:417); or
- (g)(e) the full-length coding sequence of the cDNA deposited under ATCC accession number 203115;

 wherein said polypeptide induces proliferation of stimulated T lymphocytes in a mixed lymphocyte reaction.
- 121. (Currently amended) An isolated nucleic acid of Claim 119 having at least 90% nucleic acid sequence identity to:
- (a) a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418);
- (b) a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418), lacking its associated signal peptide;
- (c) a nucleic acid sequence encoding the extracellular domain of the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418);
- (d) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in Figure 300 (SEQ ID NO:418), lacking its associated signal peptide;
- (e) the nucleic acid sequence of SEQ ID NO: 417 shown in Figure 299 (SEQ ID NO:417);
- (f)(d) the full-length coding sequence of the nucleic acid sequence of SEQ ID NO: 417 shown in Figure 299 (SEQ ID NO:417); or
- (g)(e) the full-length coding sequence of the cDNA deposited under ATCC accession number 203115;
 - wherein said polypeptide induces proliferation of stimulated T lymphocytes in a mixed lymphocyte reaction.

- 122. (Currently amended) An isolated nucleic acid of Claim 119 having at least 95% nucleic acid sequence identity to:
- (a) a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418);
- (b) a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418), lacking its associated signal peptide;
- (c) a nucleic acid sequence encoding the extracellular domain of the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418));
- (d) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in Figure 300 (SEQ ID NO:418), lacking its associated signal peptide;
- (e) the nucleic acid sequence of SEQ ID NO: 417 shown in Figure 299 (SEQ ID NO:417);
- (f)(d) the full-length coding sequence of the nucleic acid sequence of SEQ ID NO: 417 shown in Figure 299 (SEQ ID NO: 417); or
- (g)(e) the full-length coding sequence of the cDNA deposited under ATCC accession number 203115;

 wherein said polypeptide induces proliferation of stimulated T lymphocytes in a mixed lymphocyte reaction.
- 123. (Currently amended) An isolated nucleic acid of Claim 119 having at least 99% nucleic acid sequence identity to:
- (a) a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418);
- (b) a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418), lacking its associated signal peptide;
- (c) a nucleic acid sequence encoding the extracellular domain of the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418);
- (d) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in Figure 300 (SEQ ID NO:418), lacking its associated signal peptide;

- (e) the nucleic acid sequence of SEQ ID NO: 417 shown in Figure 299 (SEQ ID NO:417);
- (f)(d) the full-length coding sequence of the nucleic acid sequence of SEQ ID NO: 417 shown in Figure 299 (SEQ ID NO:417); or
- (g)(e) the full-length coding sequence of the cDNA deposited under ATCC accession number 203115;

 wherein said polypeptide induces proliferation of stimulated T lymphocytes in a mixed lymphocyte reaction.
- 124. (Currently amended) An isolated nucleic acid comprising:
- (a) a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418);
- (b) a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418), lacking its associated signal peptide;
- (c) a nucleic acid sequence encoding the extracellular domain of the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418);
- (d) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in Figure 300 (SEQ ID NO:418), lacking its associated signal peptide;
- (e) the nucleic acid sequence of SEQ ID NO: 417 shown in Figure 299 (SEQ ID NO:417);
- (f)(d) the full-length coding sequence of the nucleic acid sequence of SEQ ID NO: 417 shown in Figure 299 (SEQ ID NO:417); or
- (g)(e) the full-length coding sequence of the cDNA deposited under ATCC accession number 203115.
- 125. (Currently amended) The isolated nucleic acid of Claim 124 comprising a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO: 418).

- 126. (Currently amended) The isolated nucleic acid of Claim 124 comprising a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO: 418), lacking its associated signal peptide.
- 127. (Currently amended)The isolated nucleic acid of Claim 124 comprising the nucleic acid sequence encoding the extracellular domain of the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418), lacking its associated signal peptide.
- 128. Canceled.
- 129. (Currently amended) The isolated nucleic acid of Claim 124 comprising the nucleic acid sequence of SEQ ID NO: 417 shown in Figure 299 (SEQ ID NO:417).
- 130. (Currently amended) The isolated nucleic acid of Claim 124 comprising the full-length coding sequence of the nucleic acid sequence of SEQ ID NO: 417 shown in Figure 299 (SEQ ID NO:417).
- 131. (Previously presented) The isolated nucleic acid of Claim 124 comprising the full-length coding sequence of the cDNA deposited under ATCC accession number 203115.
- 132. (Currently amended) An isolated nucleic acid that hybridizes under stringent conditions to:
- (a) a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO: 418);
- (b) a nucleic acid sequence encoding the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO: 418), lacking its associated signal peptide;
- (c) a nucleic acid sequence encoding the extracellular domain of the polypeptide of SEQ ID NO: 418 shown in Figure 300 (SEQ ID NO:418);

- (d) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in Figure 300 (SEQ ID NO:418), lacking its associated signal peptide;
- (e) the nucleic acid sequence of SEQ ID NO: 417 shown in Figure 299 (SEQ ID NO:417);
- (f)(e) the full-length coding sequence of the nucleic acid sequence of SEQ ID NO: 417 shown in Figure 299 (SEQ ID NO:417); or
- (g)(f) the full-length coding sequence of the cDNA deposited under ATCC accession number 203115;

 wherein said stringent conditions employ hybridization using 50% formamide, 5X SSC, 50 mM sodium phosphate (pH 6.8), 0.1% sodium pyrophosphate, 5X Denhardt's solution, sonicated salmon sperm DNA (50 μg/ml), 0.1% SDS, and 10% dextran sulfate at 42°C, and washes at 42°C in 0.2X SSC, at 55°C in 50% formamide followed by a high-stringency wash at 55°C in 0.1X SSC, EDTA.
- 133. Canceled.
- 134. (Previously presented) The isolated nucleic acid of Claim 52 which is at least 10 nucleotides in length.
- 135. (Currently amended) A vector comprising the nucleic acid of Claim 124 119.
- 136. (Previously presented) The vector of Claim 135, wherein said nucleic acid is operably linked to control sequences recognized by a host cell transformed with the vector.
- 137. (Previously presented) A host cell comprising the vector of Claim 135.
- 138. (Previously presented) The host cell of Claim 137, wherein said cell is a CHO cell, an *E. coli* or a yeast cell.